

**EXHIBIT A**

1: CAA09617. serum albumin [Mu...  
[gi:3647327]

BLink, Domains, Nucleotide, Related Sequences, Domain Relatives,  
Taxonomy, LinkOut

LOCUS CAA09617 608 aa linear ROD 02-FEB-2000  
 DEFINITION serum albumin [Mus musculus].  
 ACCESSION CAA09617  
 VERSION CAA09617.1 GI:3647327  
 DBSOURCE embl locus MMU011413, accession [AJ011413.1](#)  
 KEYWORDS .  
 SOURCE house mouse.  
 ORGANISM Mus musculus  
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
 Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.  
 REFERENCE 1 (residues 1 to 608)  
 AUTHORS Cable,J., Harris,P.D., Tinsley,R.C. and Lazarus,C.M.  
 TITLE Phylogenetic analysis of Gyrodactylus spp. (Platyhelminthes:  
 Monogenea) using rDNA sequences  
 JOURNAL Can. J. Zool. 77, 1439-1449 (1999)  
 REFERENCE 2 (residues 1 to 608)  
 AUTHORS Van Reeth,T., Gabant,P., Dreze,P., Szpirer,J. and Szpirer,C.  
 JOURNAL Unpublished  
 REFERENCE 3 (residues 1 to 608)  
 AUTHORS Van Reeth,T.  
 TITLE Direct Submission  
 JOURNAL Submitted (18-SEP-1998) Van Reeth T., Departement de Biologie  
 Moleculaire, Universite Libre de Bruxelles, Laboratoire de Biologie  
 du Developpement, rue des chevaux, 67, 1640 Rhode St Genese,  
 BELGIUM  
 FEATURES Location/Qualifiers  
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 Protein 1..608  
 CDS 1..608  
 /coded\_by="AJ011413.1:34..1860"  
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 ORIGIN  
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 61 dehaklvqev tdfaktcvad esaancdksl htlfgdklca ipnlrenyge ladccatkqep  
 121 ernecflqhk ddnpslppfe rpeaeamcts fkennpttfmg hylhevarrh pyfyapelly  
 181 yaeqyneilt qcceadkes cltpkldgvk ekalvssvrq rmkcsmqkf gerafkawav  
 241 arlsqtfpna dfaeitklat dltnvnkecc hgdllcadd raelakymce nqatissklq  
 301 tccdkplkk ahclsevehd tmpadlpaia adfvvedqevc knyaekdvf lgtflyeysr  
 361 rhpdysvsll lrlakkyeat lekccaeap pacygtvlae fqplveepkn lvktncdlye  
 421 klgeygfqna ilvrytqkap qvstptlvea arnlgrvgtk cctlpedqrl pcvedylsai  
 481 lnrvcllhek tpvsehvtkc csgslverrp cfsaltvdet yvpkefkaet ftfhsdiclt  
 541 pekekqikkq talaelvkhk pkataeqlkt vmddfaqfld tcckaadkdt cfstegpnlv  
 601 trckdala

The screenshot shows the NCBI Protein search interface. At the top, there's a navigation bar with links to PubMed, Nucleotide, Protein, Genome, Structure, PopSet, Taxonomy, OMIM, and Book. Below the navigation bar, a search bar displays "Search [Protein] for CAA09617". To the right of the search bar are "Go" and "Clear" buttons. Below the search bar, there are buttons for "Limits", "Preview/Index", "History", "Clipboard", and "Details". Under "Clipboard", there are buttons for "Display" (set to "default"), "Save" (as "Text"), and "Add to Clipboard".

1: CAA23753. reading frame  
HSA...[gi:28590]

BLink, Domains, Nucleotide, OMIM, Related Sequences, Domain Relatives,  
PubMed, SNP, Taxonomy, LinkOut

LOCUS CAA23753 609 aa linear PRI 12-SEP-1993  
 DEFINITION reading frame HSA [Homo sapiens].  
 ACCESSION CAA23753  
 VERSION CAA23753.1 GI:28590  
 DBSOURCE embl locus HSALB1, accession V00494.1  
 KEYWORDS .  
 SOURCE human.  
 ORGANISM Homo sapiens Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 REFERENCE 1 (residues 1 to 609)  
 AUTHORS Dugaiczyk,A., Law,S.W. and Dennison,O.E.  
 TITLE Nucleotide sequence and the encoded amino acids of human serum albumin mRNA  
 JOURNAL Proc. Natl. Acad. Sci. U.S.A. 79 (1), 71-75 (1982)  
 MEDLINE 82105994  
 PUBMED 6275391  
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           /db\_xref="SWISS-PROT:P02768"  
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 121 grnecflqhk ddnpnlprlv rpevdvmcta fhdneetflk kylyeiarrh pyfyapellf  
 181 fakrykaft eccqaadkaa cilpkldelr degkassakq rlkcaslqkf gerafkawav  
 241 arlsqrfpka efaevsklv t dlkvhtecc hgdllcadd radlakyice nqdsissklk  
 301 eccekpllek shciaevend empadlpsla adfveskdvc knyaekdvf lgmflyeyar  
 361 rhpdysvvll lrlaktyett lekccaaadp hecyakvfde fkplveepqn likqncelfe  
 421 qlgeykfqna llvrytkvp evstptlvev srnlkgvgsk cckhpeakrm pcaedylsvv  
 481 lnqlcvlhek tpvsdrvtkc cteslvnrrp cfsalevdet yvpkefnetaet ftfhadictl  
 541 sekerqikkq talvelvkhk pkatkeqlka vmddfaafve kcckaddket cfaeegkkvl  
 601 aasqaalgl  
 //

Revised: October 24, 2001.

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